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Sequence 36, Appl
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Sequence 249, App
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Sequence 25, Appl
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| ### ### ############################## | SEGURACE 1. ADDITICATION US/09704611 Parent No. (554284 GENERAL INFORMATION: APPLICANT: USCREPARATE TITLE OF INVENTION: APPLICANT: USCREPARATE TORREST FILING DATE: 2000-11-03 SEQ ID NO 15 SE | / 353 ILGSERKKTIANYLVWRMVYSRIPNLSRRFQYRWLEFSRVIQGTTTLLPQWDKCVNFIES 412 |

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; Sequence 2, Application US/08289112
; Patent No. 5688640
; GENERAL INFORMATION:
FILING DATE: 11-NOV-1994
INFORMATION FOR SEQ ID NO: 36:
SEQUENCE CHARACTERISTICS:
                                                            : 753 amino acids
amino acid
                                                                                                                       MOLECULE TYPE: protein
                                                                                                     TOPOLOGY:
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                                                                                                   SVFIRLYVSPDDKASNEHILKLDQATLSLAVREDYLDNSTEAKSYRDALYKFMVDTAVLL 260
                                                                                                                           261 GANSSRA-EHDMKSVLRLEIKIAEIMIPHENRTSEAM-YNKMNISELSAMIPQFDWLGYI 318
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                    SCMNEKAIEKADAKPLLHILRHSPFRWPVLESNIGPEGVWSERKFSLLQTLATFRGQYSN 200
                                                        117 ACMNETRIEELRAKPLMELIERLG-GWNI-----TGPWAKDNFQ--DILQVVTAHYRT 166
                                                                                                                                                                                                                            226 GGGDEEAIRPQMQQILDFETALANITIPQEKRRDEELIYHKVTAAELQTLAPAINWLPF- 284
                                                                                                                                                                                                                                                                     319 KKVIDTRLYPHLKDISPSENVVVRVPQYFKDLFRILGSERKKTIANYLVWRMVYSRIPNL 378
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CLASTFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP94/03706
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APPLICATION NUMBER: US/08/646,273
FILING DATE: 16-NOY-1994
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Patent No. 6066502
GENERAL INFORMATION:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
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                                                                                                                                                                                                    21 IALVVFVGGTLVLGTILFLVSQGLLSLQAKQEYCLKPECIEAAAAILSKVNLSVDPCDNF
/ Match 153, Length 753, Local Similarity 37.6%; Pred. No. 8.3e-115, 1ength 753, 1es 281; Conservative 126, Mismatches 276; Indels 64
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TITLE OF INVENTION: Endothelin Converting Enzyme-1: A
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725 EEFQKAFNCPPNSTMNRGMDSCRLW 749
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FILING DATE: 16-NOV-1994
CLASSIPICATION: 435
PRIOR APPLICATION DATE:
APPLICATION NUMBER: PCT/EP94/03706
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1101 Connecticut Avenue
                                                                                                                                                                                                                                                                                                                                                                                            US-08-646-273-23; Sequence 23, Application US/08646273; Patent No. 6066502; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 11-NOV-1994
INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
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TYPE: amino acid
O'DPOLOGY: 1:-
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MEDIUM TYPE: Diskett
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USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         203 FIRLYVSPDDKASNEHILKLDQATLSLAVREDYLDNSTEAKSYRDALYKFMVDTAVLLGA 262
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Membrane-Bound Metalloprotease That Catalyzes The Proteolytic Activation of Big Endothelin-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 31.5%; Score 1248.5; DB 1; Length 758; Best Local Similarity 37.9%; Pred. No. 5.8e-114; Matches 282; Conservative 121; Mismatches 281; Indels 61;
                                                                                                                                                                                                                                                                     SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                               10-APR-1994
10-APR-1994
11 435
  TITLE OF INVENTION: Membrane-Bound N
TITLE OF INVENTION: Proteolytic Acti
NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
ADDRESSEE: Arnold, White & Durkee
                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                            NAME: Parker, David L. REGISTRATION NUMBER: 32,165
                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: UTS
TELECOMMUNICATION INFORMATION:
TELEPHONE: 713-787-1400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acids
TOPOLOGY: 1:-
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INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
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                                                                  ADDRESSEE: Arnold,
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                                                                                                                                                    COUNTRY: USA
ZIP: 77210-4433
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STREET: F.C.
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CLASSIFICATION:
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APPLICANT: Kroeger, Burkhard, Seulberger, Harald, Meyer, Thomas, Schmidt, Papplicant: Martin, Jacob, Elard, Otter, Rainer, Subkowski, Thomas, Hillen, Heinz. TITLE OF INVENTION: Endothelin Converting Enzyme (ECE).
                                                                                                                                           66 YACGGWIKANPVPDGHSRWGTFSNLWEHNQAIIKHLLENSTA--SVSEAERKDQEYYRAC 123
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516 DL-----YFENAMRF----FNFSWRVTADQLRKAPNRDQWSMTPPMVNAYYSPT 560
                                                                                   545 TNQIRFPAGELQKPFFWGTEYPRSLSYGAIGVIVGHEFTHGFDNNGRKYDKNGNLDPWWS 604
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APPLICANT: WEL, Ming-Hui et al.

TITLE OF INVENTION: ISOLATED HUMAN ENZYME PROTEINS, NUCLEIC

TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN ENZYME PROTEINS, AND USES

TITLE OF INVENTION: THEREOF

FILE REFERENCE: CLO01200

CURRENT APPLICATION NUMBER: US/09/819,989

CURRENT FILING DATE: 2001-03-29

NUMBER OF SEQ ID NOS: 4

SOFTWARE: FastSEQ for Mindows Version 4.0
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                                                                                                             Query Match 31.4%; Score 1241.5; DB 3; Length 754; Best Local Similarity 37.9%; Pred. No. 2.8e-113; Matches 282; Conservative 120; Mismatches 282; Indels 61; Gaps
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; Patent No. 6482629
; GENERAL INFORMATION:
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CORRESPONDENCES: 36
CORRESPONDENCE ADDRESS:
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PRIOR APPLICATION DATA:
APPLICATION NOMBER: PCT/EP94/03706
FILING DATE: 11-NOV-1994
INFORMATION FOR SGO ID NO: 30:
SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER: US/08/646,273
FILING DATE: 16-NOV-1994
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1101 Connecticut Avenue
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MEDIUM TYPE: Diskett
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Sequence 2, Application US/09819989
Patent No. 6482629
GENERAL INFORMATION:
APPLICANT: WEL, Ming-Hui et al.
TITLE OF INVENTION: ISOLATED HUMAN ENZYME PROTEINS, NUCLEIC
TITLE OF INVENTION: ACID MOLECULES ENCODING HUMAN ENZYME PROTEINS, AND USES
TITLE OF INVENTION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          173 NIGP---EGVWSERKFSLLQTLATFRGQYSNSVFIRLYVSPDDKASNEHILKLDQATLSL 229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                230 AVREDYLDNSTEAKSYRDALYKFMVDTAVLLGANSSRAEHDMKSVLRLEIKIAEIMIPH- 288
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         318 ORRDEEKIYHXMSISELQALAPSMDWLEFLSFLLSP-----LELSDSEPVVVYGMDYLQ 371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          349 DLFRILGSERKKTIANYLVWRMYSRIPNLSRRFQYRWLEFSRVIQGT-TTLLPQWDKCV 407
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            53 YCLKPECIEAAAAILSKVNLSVDPCDNFFRFACDGWISNNPIPEDMPSYGVYPWLRHNVD 112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               93 TCLTEACIRVAGKILESLDRGVSPCEDFYQFSCGGWIRRNPLPDGRSRWNTFNSLWDQNQ 152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   201 KIGGWNITGPWDQDNF--MEVLKAVAGTYRATPFFTVYISADSKSSNSNVIQVDQSGLFL 258
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       4 ETGSS---VETG-KKANR---GTRIAL-VVFVGGTLVLGTILF--LVSQGL-LSLQAKQE 52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 30.9%; Score 1222.5; DB 4; Length 765; Best Local Similarity 35.4%; Pred. No. 2.2e-111; Matches 270; Conservative 145; Mismatches 302; Indels 45; Gaps
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                                                                                        TYPE: PRT
ORGANISM: Human
SEQ ID NO 4
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211 NSSSEAEQKTQRFYLSCLQVERIEELGAQPLRDLI------EKIGGWNITGPWDQ 259
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                                                                                                                                                                                                                                                                                                                                                                                 92 QKGTRQLLGSRTQLELVLAGASLLLAALLLGCLVALGVQYHRDPSHSTCLTEACIRVAGK 151
                                                                                                                                                                                                                                                                                                                                                                                                                                                         66 ILSKVNLSVDPCDNFFRFACDGWISNNPIPEDMPSYGVYPWLRHNVDLKLKELLEKSISR 125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        152 ILESLDRGVSPCEDFYQFSCGGWIRRNPLPDGRSRWNTFNSLWDQNQAILKHLLENT-TF 210
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           126 RRDTEAIQKAKILYSSCMNEKAIEKADAKPLLHILRHSPFRWPVLESNIGP----EGVWSE 182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        183 RKFSLLQTLATFRGQYSNSVFIRLYVSPDDKASNEHILKLDQATLSLAVREDYLDNSTEA 242
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        260 DNF--MEVLKAVAGTYRATPFFTVYISADSKSSNSNVIQVDQSGLFLPSRDYYL-NRTAN 316
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          243 KSYRDALYKFMVDTAVLLGANSSRAEHDMKSVLRLEIKIAEIMIPH-ENRTSEAMYNKMN 301
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             302 ISELSAMIPQFDWLGYIKKVIDTRLYPHLKDISPSENVVVRVPQYFKDLFRILGSERKKT 361
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            377 İSELQALAPSMDWLEFLSFLLSP-----LELSDSEPVVVYGMDYLQQVSELINRTEPSI 430
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            362 IANYLVWRMVYSRIPNLSRRFQYRWLEFSRVIQGT-TTLLPQWDKCVNFIESALPYVVGK 420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 421 MFVDVYFQEDKKEMMEELVEGVRWAFIDMLEKENEWMDAGTKRKAKEKARAVLAKVGYPE 480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 481 FIMNDTHVNEDLKAIKFSEADYFGNVLQTRKYLAQSDFFWLRKAVPKTEWFTNPTTVNAF 540
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 541 YSASTNQIRFPAGELQKPFFWGTEYPRSLSYGAIGVIVGHEFTHGFDNNGRKYDKNGNLD 600
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                                                                                                                                                                                                                                                                                                                                                      10 ETCKKANRGTRIAL-VVFVGGTLVLGTILF--LVSQGL-LSLQAKQEYCLKPECIEAAAA 65
                                                                                                                                                                                                                                                    Query Match 30.8%; Score 1221; DB 4; Length 811; Best Local Similarity 35.2%; Pred. No. 3.4e-111; Matches 264; Conservative 144; Mismatches 303; Indels 38; Gaps
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CURRENT APPLICATION NUMBER: US/09/819,989 CURRENT FILING DATE: 2001-03-29 NUMBER OF SEQ ID NOS: 4 SOFTWARE: PARLSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 4, Application US/09667373; Patent No. 6524840; GENERAL INFORMATION: APPLICANT: Donoho, Gregory
                                                                                                                                                                                  ORGANISM: Human
                                                                                                       SEQ ID NO 2
LENGTH: 811
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-667-373-4
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              APPLICANT: Nehls, Michael C.
APPLICANT: Friedrich, Glenn
APPLICANT: Zandrowicz, Brian
APPLICANT: Sands, Arthur T.
TITLE OF INVENTION: No. 6524840el Human Endothelin Converting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  164 OKGTROLLGSRIOLELVLAGASLLLAALLLGCLVALGVQYHRDPSHSTCLTEACIRVAGK 223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             66 ILSKVNLSVDPCDNFFRFACDGWISNNPIPEDMPSYGVYPWLRHNVDLKLKELLEKSISR 125
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      126 RRDTEAIQKAKILYSSCMNEKAIEKADAKPLLHILRHSPFRWPVLESNIGP---EGVWSE 182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               283 NSSSEAEOKTORFYLSCLQVERİEELGAQPLRDLI-------EKIGGWNITGPWDQ 331
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 243 KSYRDALYKFMVDTAVLLGANSSRAEHDMKSVLRLEIKIAEIMIPH-ENRTSEAMYNKAN 301
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              38; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
30.8%; Score 1221; DB 4; Length 883;
Best Local Similarity 35.2%; Pred. No. 3.9e-111;
Matches 264; Conservative 144; Mismatches 303; Indels 38
                                                                                                                                                     FILE REFERENCE: LEX-0043-USA
CURRENT APPLICATION NUMBER: US/09/667,373
CURRENT FILING DATE: 2000-09-22
PRIOR APPLICATION NUMBER: US 60/156,102
                                                                                                                                                                                                                                         PRIOR FILING DATE: 1999-09-24
PRIOR APPLICATION NUMBER: US 60/176,689
BYROR FILING DATE: 2000-01-18
NUMBER OF SEQ ID NOS: 5
SOFTWARE: FastSEQ for Windows Version 4.0
Turner, C. Alexander Jr.
                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: homo sapiens
                                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 4
LENGTH: 883
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APPLICANT:
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61 EAAAAILSKVNLSVDPCDNFFRFACDGWISNNPIPEDMPSYGVYPWLRHNVDLKLKELLE 120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      231 GPWDQDNF--MEVLKAVAGTYRATPFFTVYVSADSKSSNSNIIQVDQSGLFLPSRDYYL- 287
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Matches 265; Conservative 138; Mismatches 313; Indels 38; Gaps
                                                                                                                                                                                                                                       APPLICANT: Yanagisawa, Masashi
TITLE OF INVENTION: RECOMBINANT ENDOTHELIN CONVERTING
TITLE OF INVENTION: ENZYME-2 AND ITS USE IN ECE INHIBITOR SCREENING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29.9%; Score 1184; DB 1; Length 787; 35.1%; Pred. No. 1.4e-107;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
721 ISNSEEFQKAFNCPPNSTMNRGMDSCRLW 749
                                              856 LŚNŚRDFLRHFGĆPVGSPMNPG-QLĆEVW 883
                                                                                                                                                                                                                                                                                                                                                                                                                                    STAID.
COUNTY: USA
ZIP: 77210-4433
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
"""THER: IBM PC compatible
"""THER: """""""" PC "OOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/08/574,763
FILING DATE: Concurrently herewith
CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Arnold, White & Durkee STREET: P. O. Box 4433 CITY: Houston
                                                                                                                                                                   Sequence 2, Application US/08574763
Patent No. 5736376
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Corder, Timothy S. REGISTRATION NUMBER: 38,414
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: UT: TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (512) 418-3000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (512) 474-7577
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
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                                                                                                                  RESULT 11
US-08-574-763-2
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106 WLRHNVDLKLKELLEKSISRRRDTEAlQKAKILYSSCMNEKALEKADAKPLLHILRHSPF 165
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       718 NGAISNSEEFQKAFNCPPNSTMNRGMDSCRLW 749
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 745 LGSVSQFEEFGRAFHCPKDSPMNPA-HKCSVW 775
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP94/03706
FILING DATE: 11-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Wordberfect version 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/646,273
FILING DATE: 16-NOV-1994
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             3: Keil & Weinkauf
1101 Connecticut Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; Sequence 19, Application US/08646273
; Patent No. 6066502
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     : 567 amino acids
amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                231 LLYKAQGVYSAAALFSLTVSLDDRNSSRYVIRIDQDGLTLPERTLYLAQDEDSEKILAAY 290
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            80 FFRFACDGWISNNPIPEDMPSYGVYPWLRHNVDLKLKELLEKSISRRRDTEAIQKAKILY 139
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      140 SSCMNEKALEKADAKPLLHILRHSPFRWPVLESNIGPEGVW----SER----KFSLLQ 189
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                                                                                                                                                                                                                                                                                                                                                                                                  640 EGNLRPWWQNESLAAFRNHTACIEEQYSQY--QVNGEKINGRQTLGENIADNGGLKAAYN 697
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                                                                                                                                                                                      476 VGYPEFIMNDTHVNEDLKAIKFSEADYFGNVLQTRKYLAQSDFFWLRKAVPKTEWFTNPT 535
                                                                                                                                                                                                                        521 IGFPDFILEPKELDDVYDGYBVSEDSFFQNMLNLYNFSAKVMADQLRKPPSRDQWSMTPQ 580
                                                                                                                                                                                                                                                                             536 TVNAFYSASTNQIRFPAGELQKPFFWGTEYPRSLSYGAIGVIVGHEFTHGFDNNGRKYDK 595
                                                                                                                                                                                                                                                                                                                            581 TVNAYYLPTKNEIVFPAGILQAPFY-TCNHPQALNFGGIGVVMGHELTHAFDDQGREYDK 639
                                                                                                                                                                                                                                                                                                                                                                       596 NGNLDPWWSTESEEKFKEKTKCMINQYSNYYWKKAGINVKGKRTLGENIADNGGLREAFR 655
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   656 AYRKWINDRROGLEEPLLPGITFINNQLFFLSYAHVRCNSYRPEAAREQVQIGAHSPPQF 715
357 ERKKTIANYLVWRMVYSRIPNLSRRFQYRWLEFSRVIQGT-TTLLPQWDKCVNFIESALP 415
                                          402 TEPSVLNNYLIWNLVQKTTSSLDHRFESAQEKLLETLYGTKKSCTPRWQTCISNTDDALG 461
                                                                                       416 YVVGKMFVDVYFQEDKKEMMEELVEGVRWAFIDMLEKENEWMDAGTKRKAKEKARAVLAK 475
                                                                                                                                    462 FALGSLFVKATFDRQSKEIAEGMISEIRVÄFEBAL-GHLVWMDEKTRQAAKEKADAIYDM 520
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24.8%; Score 983.5; DB 3; Length 775;
Best Local Similarity 31.0%; Pred. No. 8.9e-88;
Matches 233; Conservative 158; Mismatches 294; Indels 67;
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TITLE OF INVENTION: No. 6255468el Compounds
FILE REFERENCE: GP30124
CURRENT APPLICATION NUMBER: US/09/305,640B
CURRENT FILING DATE: 1999-05-05
UNDRENT FILING DATE: 1999-05-05
NUMBER OF SEQ ID NOS: 4
SOFTWARE: FASTSEQ for Windows Version 3.0
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Patent No. 6255468
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ORGANISM: Homo sapiens
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APPLICANT: Kroeger, Burkhard, Seulberger, Harald, Meyer, Thomas, Schmidt, APPLICANT: Martin, Jacob, Elard, Otter, Rainer, Subkowski, Thomas, Hillen, Heinz. TITLE OF INVENTION: Endothelin Converting Enzyme (ECE).

NUMBER OF SEQUENCES: 36

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                           540 FYSASTNGIRFPAGELOKPFFWGTEYPRSLSYGALGVIVGHEFTHGFDNNGRKYDKNGNL 599
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                                                                                                              420 KMFVDVYFQEDKKEMMEELVEGVRWAFIDMLEKENEWMDAGTKRKAKEKARAVLAKVGYP 479
                                                                                                                                                                                                                            480 EFIMNDTHVNEDLKAIKFSEADYFGNVLQTRKYLAQSDFFWLRKAVPKTEWFTNPTTVNA 539
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349 KITPHLRWKWLLDQIFQ------EDFSEEEEVVLLATDYMQQVSQLIRSTPHRVLHNYL 401
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24.3%; Score 963.5; DB 3; Length 567;
Best Local Similarity 37.7%; Pred. No. 4.9e-86;
Matches 226; Conservative 96; Mismatches 216; Indels 61; Gaps
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APPLICANT: Kroeger, Burkhard, Seulberger, Harald, Meyer, Thomas, Schmidt, APPLICANT: Martin, Jacob, Elard, Otter, Rainer, Subkowski, Thomas, Hillen, Heinz TITLE OF INVENTION: Endothelin Converting Enzyme (ECE).

CORRESPONDENCE: 36

CORRESPONDENCE ADDRESS:
                                         305 LSAMIPQFDWLGY-----IKKVIDTRLYPHLKDISPSENVVVRVPQYFKDLFRILGSERK 359
                                                                                                                                                                                                                                                                 360 KTIANYLVWRMVYSRIPNLSRRFQYRWLEF-SRVIQGTTTLLPQWDKCVNFIESALPYVV 418
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246 RDALYKFMVDTAVLLGANSSRAEHDMKSVLRLEIKIAEIMIPH-ENRTSEAMYNKMNISE 304
                                                                                                                                                                                               269 LYKLTPAFDWQGYLTATELKGKIDT-------IQVNQPDYFQGLNTIIQNTPL 314
                                                                                                                                                                                                                                                                                                                            315 DIWKAYFKLHLINNFSPLLNSAFVDNKFDFYNKTLRDITEQEARWKRGVQLVDKVLGDGI 374
                                                                                                                                                                                                                                                                                                                                                                                                                                                                375 GKLYVEKYFPAEKKQQMELLVQNLIRAY-DQSITELDWMSPATKIQARKKLSHMSIKIGY 433
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COMPUTER: Diskette, 3.5 inch, 1.4 Mb storage
COMPUTER: IBM AT-compatible, 80486 processor
ODRPATING SYSTEM: MS-DOS version 6.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          719 GAISNSEEFQKAFN-----CPPNSTMN 740
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CLASSIFICATION: 435
PLASSIFICATION DATA:
APPLICATION DATA:
FILING DATE: 11-NOV-1994
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
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FILING DATE: 16-NOV-1994
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Patent No. 6066502
GENERAL INFORMATION:
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amino acid
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TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
TITLE OF INVENTION: BAUMANNII FOR DIAGNOSTICS AND THERAPEUTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WDKÇVNFIESALPYVVGKMFVDVYFQEDKKEMMEELVEGVRWAFIDMLEKENEWMDAGTK 462
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                                                                  166 RWPVLESNIGPEGVWSERKFSLLQTLATFRGQYSNSVFIRLYVSPDDKASNEHILKLDQA 225
                                                                                                                         67 GWNI-----TGPWDKDNFQ--DTLOVVTSHYHTSPFFSVYVSADSKNSNSNVIQVDQS 117
                                                                                                                                                                                                                                                          118 GLGLPSRDYYL-NKTENEKVLTGYLNYMVOLGKLLGGGAEDTIRPOMOQILDFETALANI 176
                                                                                                                                                                                                                                                                                                                        285 MIPHENRISEAM-YNKANISELSAMIPQFDWLGYIKKVIDTRLYPHLKDISPSENVVVRV 343
                                                                                                                                                                                                                                                                                                                                                                                          177 TIPQEKRRDEELIYHKVTAAELQTLAPAINWLPF----LNTIFYP--VEINESEPIVIYD 230
                                                                                                                                                                                                                                                                                                                                                                                                                                                             344 PQYFKDLFRILGSERKKTIANYLVWRMVYSRIPNLSRRFQYRWLEFSRVIQGT-TTLLPO 402
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           231 KEYLSKVSTLINSTDKCLLNNYMIWNLVRKTSSFLDQRFQDADEKFMEVMYGTKKTCLPR 290
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         350 KSAKEKADAIYNMIGYPNFIMDPKELDKVFNDYTAVPDL-----YFENAMRF---- 396
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     568 SLSYGAIGVIVGHEFTHGFDNNGRKYDKNGNLDPWWSTESEEKFKEKTKCMINQYSNYYW 627
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  628 KKAGLNVKGKRTLGENIADNGGLREAFRAYRKWINDRRQGLEEPLLPGITFTUNQLFFL 686
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12 W-EHNQAI-IKHLLENSTA--SVSEAERKDQEYYRACMNETRIEELKAKPLMELIEKLG- 66
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20.0%; Score 790; DB 4; Length 690;
Best Local Similarity 30.3%; Pred. No. 9.4e-69;
Matches 209; Conservative 132; Mismatches 279; Indels 70; Gaps
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CURRENT APPLICATION NUMBER: US/09/328,352
CURRENT FILING DAIE: 1999-06-04
NUMBER OF SEQ ID NOS: 8252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 5182, Application US/09328352 Patent No. 6562958 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Acinetobacter baumannii
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LENGTH: 690
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Query Match
Best Local Similarity 51.8%; Pred. No. 8.5e-44;
Best Local Similarity 51.8%; Pred. No. 8.5e-44;
Matches 101; Conservative 32; Mismatches 55; Indels 7; Gaps 5;

y 555 LQXPFWGTEYPRSLSYGAIGVIVGHEFTHGFDNNGRKYDKNGNLDPWNSTESEKFKEK 614

y 615 TKCMINQYSNYYWKKAGLIVYGHETHAFDDGGREYDKDGNLRPWWRNSSVBAFKQO 60

y 615 TKCMINQYSNYYWKKAGLIVYGRATLGENIADNGGLREAPRAYRWINDRRQGEEPLLP 674

b 61 TACMYEQYGNY -- SYNGEPVNGRHILGENIADNGGLKAAYRAYQNWV -- KKNGAEQ-TLP 115

y 675 GITFTNNQLFFLSYAHVRCNSYRPEAAREQVQIGAHSPPOFRVNGAISNSEBFQKAFNCP 734

y 735 PNSTMNRGMDSGRLW 749
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earch completed: December 3, 2003, 07:31:49 ob time : 32.3726 secs

176 PGSPMN-PHHKCEVW 189

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                                                                                                                                                     December 3, 2003, 07:21:37; Search time 497.375 Seconds (without alignments) 280.074 Million cell updates/sec
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Sequence 526, 1
Sequence 526, 1
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Sequence 2, 1
Sequence 6, 1
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1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/BCT_NEW_PUB.pep:*

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GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd.
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US-00-10-147-928-4

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US-10-1017-273A-6

US-10-1017-273A-5

US-00-978-298-5

US-00-978-298-5

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US-00-99-825A-526

US-00-978-192A-526

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Listing first 45 summaries

    protein search, using sw model

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Maximum DB seq length: 2000000000
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Match Length DB
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APPLICANT: SOLVAY PHARMACEUTICALS B.V.
TITLE OF INVENTION: NO. US20031080877Ael Human Enzymes of the Metalloprotease Famil-
FILE REPERENCE: SPW 99.09 /H 99.26-WO
CURRENT APPLICATION NUMBER: US/10/147,928
CURRENT FILING DATE: 2002-05-20
NUMBER OF SEQ ID NOS: 31
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US-09-978-423A-526
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US-10-134-526
US-10-134-526
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; Publication No. US20030180877A1
; GENERAL INFORMATION:
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SOFTWARE: FASTSEQ for Windows Version 4.0
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ORGANISM: Homo sapiens
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TITLE OF INVENTION: No. US20030180877A1el Human Enzymes of the Metalloprotease Family
FILE REFERENCE: SPW 99.09 /H 99.26-WO
CURRENT PELLICATION UMBER: US/10/147,928
CURRENT FILING DATION UMBER: US/10/147,928
NUMBER OF SEQ ID NOS: 31
                                                                                                                                                                                                                                                                                            301 DVIALYHRMGLEELQSQFGLKGFNWTLFIQTVLSS-----VKIKLLPDEEVVVYGIPVLQ 355
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     239 STEAKSYRDALYKFMVDTAVLL - GANSSR----AEHDMKSVLRLEIKIAEIMIPHENRT 292
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      356 NLENIIDTYSARTIONYLVWRLVLDRIGSLSQRFKDTRVNYRKALFGTMVEEVRWRECVG 415
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                                                                                                          242 GSNRK-VREAYLQFMVSVATLLREDANLPRDSCLVQEDMMQVLELETQLAKATVPQEERH 300
                                                                                                                                                                                                                           293 SE-AMYNKMNISELSAM--IPQFDWLGYIKKVIDTRLYPHLK-DISPSENVVVRVPQYFK 348
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Publication No. US20030180877A1
GENERAL INFORMATION:
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194 EVVGGWPVAMDR-----WNETVGLEWELERQLALMNSQFNRRVLIDLFIWNDDQNSSRH 247
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             102 GVYPWLRHNVDLKLKELLEKSISRRRDTEAIQKAKILYSSCMNEKAIEKADAKPLLHILR 161
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     137 SIFDVLRDELEVILKAVLENSTA--KDRPAVEKARTLYRSCMNOSVIEKRGSOPLLDIL- 193
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                                                                                                                                                                                                                                                                                                                                                           Query Match 33.7%; Score 1332.5; DB 15; Length 779; Best Local Similarity 39.2%; Pred. No. 7.4e-116; Matches 284; Conservative 145; Mismatches 258; Indels 37; Gaps
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                                              NUMBER OF SEQ ID NOS: 6
SOFTWARE: PatentIn version 3.1
        PRIOR FILING DATE: 2001-06-29
                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Homo sapiens
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SIFDVLRDELEVILKAVLENSTA--KDRPAVEKARTLYRSCMNQSVIEKRGSQPLLDIL- 193
                                                                       162 HSPFRWPVLESNIGPEGVWSER---KFSLLQTLATFRGQYSNSVFIRLYVSPDDKASNEH 218
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APPLICANT: Van Der Grad, Piecer H
APPLICANT: Ozn Der Grad, Piecer H
APPLICANT: Wayman, Christopher P.
IITLE OF INVENTION: Treatment of Male Sexual Dysfunction
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CURRENT APPLICATION NUMBER: US 60/265,358

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PRIOR PLING DATE: 2001-04-06
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OR APPLICATION NUMBER: GB 0108730.3

OR FILING DATE: 20001-04-06

OR APPLICATION NUMBER: GB 0120679.6

OR FILING DATE: 2001-08-24

OR FILING DATE: 2001-07-13

OR PILING DATE: 2001-07-13

OR APPLICATION NUMBER: US 00/291,722

OR APPLICATION NUMBER: US 00/291,722

OR APPLICATION NUMBER: US 00/291,722

OR APPLICATION NUMBER: US 09/895,367
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GENERAL INFORMATION:

APPLICANT: SOLVAY PHARMACEUTICALS B.V.

IITLE OF INVENTION: No. US20030180877Alel Human Enzymes of the Metalloprotease Famil)
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Godowski, Paul J.
Grimaldi, J. Christopher
Gurney, Austin L.
Hillan, Kenneth J
                                                                  Fong, Sherman
Gao, Wei-Qiang
Gerber, Hanspeter
Gerritsen, Mary E.
Goddard, Audrey
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Williams, P. Mickey
                                                                                                                                                                                                                                                                                                                                                                                                                Timothy A.
                                                                                                                                                                                                                                                                                                                                              Paoni, Nicholas F.
Roy, Margaret Ann
Shelton, David L.
                         Ferrara, Napoleon
Filvaroff, Ellen
                                                                                                                                                                                                                                                                Kljavin, Ivar J.
                                                                                                                                                                                                                                                                                Kuo, Sophia S.
Napier, Mary A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                William I
                                                                                                                                                                                                                                                                                                                                                                                                                                       Daniel
                                                                                                                                                                                                                                                                                                                              James;
                                                                                                                                                                                                                                                                                                                                                                                                                   Stewart,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Wood,
                                                                                                           APPLICANT
                                                                                                                                                                           APPLICANT
                                                                                                                                                                                                                     APPLICANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              54 CLKPECIEAAAAILSKVNLSVDPCDNFFRFACDGWISNNPIPEDMPSYGVYPWLRHNVDL 113
                                                                                                                                                                                                                                                                                                                                                                                                                                       61 ILKAVLENSTA--KDRPAVEKARTLYRSCMNQSVIEKRGSQPLLDIL-EVVGGWPVAMDR 117
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 231 VREDYLDNSTEAKSYRDALYKFMVDTAVLL--GANSSR----AEHDMKSVLRLEIKIAEI 284
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            172 SREYYFNGGSNRK-VREAYLQFMVSVATLLREDANLPRDSCLVQEDMMQVLELETQLAKA 230
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     285 MIPHENRISE-AMYNKANISELSAM--IPQFDWLGYIKKVIDTRLYPHLK-DISPSENVV 340
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          286 VYGIPYLQNLENIIDIYSARTIQNYLVWRLVLDRIGSLSQRFKDTRVNYRKALFGTMVEE 345
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  584 NTLGENIADNGGVRQAYKAYLKWM---AEGGKDQQLPGLDLTHEQLFFINYAQVWCGSYR 640
                                                                                                                                                                                                                                                                                                                                                                                                              114 KLKELLEKSISRRRDTEAIOKAKILYSSCMNEKAIEKADAKPLLHILRHSPFRWPVLESN 173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 174 IGPEGVWSER---KFSLLQTLATFRGQYSNSVFIRLYVSPDDKASNEHILKLDQATLSLA 230
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            231 TVPQEERHDVIALÝHRMGLEELQSQFGLKGFNWTLFIQTVLSS-----VKIKLLPDEEVV 285
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          341 VRVPQYFKDLFRILGSERKKTIANYLVWRMVYSRIPNLSRRFQYRWLEFSRVIQGTTTLL 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             401 PQWDKCVNFIESALPYVVGKMFVDVYFQEDKKEMMEELVEGVRWAFIDMLEKENEWMDAG 460
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                                                                                                                                                                                                                                                                                                                                                                   1 CTTPGCVIAAARILQNMDPTTEPCDDFYQFACGGWLRRHVIPETNSRYSIFDVLRDELEV 60
                                                                                                                                                                                                                                     Query Match 33.4%; Score 1322.5; DB 12; Length 691; Best Local Similarity 39.5%; Pred. No. 5.3e-115; Matches 281; Conservative 143; Mismatches 251; Indels 37; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           698 PEAAREQVQIGAHSPPQFRVNGAISNSEEFQKAFNCPPNSTMNRGMDSCRLW 749
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PILE REFERENCE: SPW 99.09 /H 99.26-WO CURRENT APPLICATION NUMBER: US/10/147,928 NUMBER OF SEQ ID NOS: 31 SOFTWARE: PATENTN VET. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 526, Application US/09978295A Patent No. US20020156006A1 APPLICANT: Ashkenazi, Avi APPLICANT: Baker Kevin P. APPLICANT: Baker Kevin P. APPLICANT: Bosnoyers, Luc
                                                                                                                                                  TYPE: PRT
CORGANISM: Homo sapiens
US-10-147-928-2
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TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic TITLE OF INVENTION: Acids Encoding the Same FILE REFERENCE: P263091011 CURRENT APPLICATION NUMBER: US/09/978,295A CURRENT FILING DATE: 2001-10-15
                                                                                                                                                           CURRENT FILING DATE: 2011-10-15
PRIOR APPLICATION NUMBER: 09/91858
PRIOR FILING DATE: 2010-10-17
PRIOR FILING DATE: 1997-10-17
PRIOR FILING DATE: 1997-10-17
PRIOR FILING DATE: 1997-10-17
PRIOR FILING DATE: 1997-11-03
PRIOR PELICATION NUMBER: 60/065311
PRIOR PELICATION NUMBER: 60/065311
PRIOR PELING DATE: 1997-11-13
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292 TSEAMYNKMNISELSAMIPQFDWLGYIKKVIDTRLYPHLKDISPSENVVVRVPQYFKDLF 351

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591 RKYDKNGNLDPWWSTESEEKFKEKTKCMINQYSNYYWKKAGLNVKGKRTLGENIADNGGL 650
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292 DEEKIYHKMSISELQALAPSMDWLEFLSFLLSP-----LELSDSEPVVVYGMDYLQQVS 345
                                                                 352 RILGSERKKTIANYLVWRMVYSRIPNLSRRFQYRWLEFSRVIQGT-TTLLPQWDKCVNFI 410
                                                                                                                                                        411 ESALPYVVGKMFVDVYFQEDKKEMMEELVEGVRWAFIDMLEKENEWMDAGTKRKAKEKAR 470
                                                                                                                                                                                                                                            471 AVLAKVGYPEFIMNDTHVNEDLKAIKFSEADYFGNVLQTRKYLAQSDFFWLRKAVPKTEW 530
                                                                                                                                                                                                                                                                                        465 AIYDMIGEPDFILEPKELDDVYDGYEISEDSFFONMLNLYNFSAKVMADQLRKPPSRDQW 524
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                                                                                                          346 ELINRIEPSILNNYLIWNLVQKTTSSLDRRFESAQEKLLETLYGTKKSCVPRWQTCISNT 405
                                                                                                                                                                                                   406 DDALGFALGSLFVKATFDRQSKEIAEGMISEIRTAFEEAL-GQLVWMDEKTRQAAKEKAD 464
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Grimaldi, J. Christopher
Gurney, Austin L.
Hillan, Kenneth J
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Patent No. US20020169284A1
GENERAL INFORMATION:
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Shelton, David L.
Stewart, Timothy A.
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Filvaroff, Ellen
Fong, Sherman
Gao, Wei-Qiang
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Goddard, Audrey
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Kuo, Sophia S.
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Botstein, David
Desnoyers, Luc
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Paoni, Nich
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US-09-978-697-526
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PRIOR APPLICATION NUMBER: 60/06314
PRIOR PLING DATE: 1997-11-01
PRIOR PELICATION NUMBER: 60/06364
PRIOR PELICATION NUMBER: 60/07450
PRIOR PELING DATE: 1997-11-12
PRIOR PELING DATE: 1997-11-11
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| | ngth 736; dels 40; Gaps | QAKQEYCL DPSHSTCL | PSYGVYPWLRHNVDLKL 1 : SRWNTFNSLWDQNQAIL 1 | LRHSPFRWPVLESNIG : EXIG | SNEHILKLDQATLSLAVR 2 | IMIPH-ENR : : VITVPQDQRR | SENVVVRVPQYFKDLF | VIQGT-TTLLPQWDKCVNFI 4 : : : : TLYGTKKSCVPRWQTCISNT 4 | 4 4 | VPKTEW 5 :: PSRDQW 5 | PRSLSYGAIGVIVGHEFTHGFDNNG 5 :: :: | GENIADNGGL 6 | SYAHVRCNSYRPEAAREQVQIGAH 7: | |
| | 1223; DB 10; Le No. 1.3e-105; smatches 307; In | RIAL-VVFVGGTLVLGTILFLVSQGL-LSL RTQLELVLAGASLLLAALLLGCLVALGVQYHR | FRFACDGWISNNPIPEDM :: : X YQFSCGGWIRRNPLPDGR | SSCMNEKAIEKADAKPLLHI : : : : : LSCLQVERIEELGAQPLRDL | YSNSVFIRLYVSPDDKA | VLLGANSSRAEHDMKSVL | IKKVIDTRLYPHLKDIS : :: LSFLLSPLELS | LSRRFQYRWLEFSR : LDRRFESAQEKLLE | MEELVEGVRWAFIDMLEKE :: : : : AEGMISEIRTAFEEAL-GQ | KAIKFSEADYFGNVLQTRKYLAQSDFFWLRKA | PAGELQKPFFWGTEYPRSLSYGA | EEKFKEKTKCMINQYSNYYWKKAGLNVKGKRTL | TFTNNOLFFL : : GLTNHOLFFV | PNSTMNRGMDSCRLW 749 : VGSPMNPG-QLCEVW 736 |
| NUMBER: 60/085582 1998-05-15 NUMBER: 60/085700 1998-05-15 NUMBER: 60/085689 1998-05-15 NUMBER: 60/085580 1998-05-15 NUMBER: 60/085580 1998-05-15 NUMBER: 60/085580 1998-05-15 NUMBER: 60/085573 NUMBER: 60/085573 NUMBER: 60/085573 NUMBER: 60/085573 | 30.9%; Score city 35.2%; Pred. nservative 145; Mi | ETGKKANRGT : : : QKGTRQLLGS | CIEAAAILSKVNLSVDPCDNFF | EKSISRRRDTEAIQKAKILY(: : : ENT-TFNSSEAEQKTQRFYI | GVWSERKFSLLQTLATFRGC | DNSTEAKSYRDALYKFMVDTA | EAMYNKMNISELSAMIPQFDWLGY : : : : EKIYHKMSISELQALAPSMDWLEF | IANYLVWRMVYSRI - : INNYLIWNLVQKTT | ESALPYVVGKMFVDVYFQEDKKEMMEELVEGVRWAFIDMLEKENEWMDAGTKRKAKEKAR | GYPEFIMNDTHVNEDL : : : GFPDFILEPKELDDVY | TNPTTVNAFYSASTNQIRFPAGEI : : MTPQTVNAYYLPTKNEIVFPAGII | RKYDKNGNLDPWWSTESEEKFKEKT : | REAFRAYRKWINDRRQGLEEPLLPGI : : : : KAAYNAYKAWL - RKHG - EEQQLPAV | PPOFRVNGAISNSEEFQKAFNCPPNSTMNRG : : PARFRVLGTLSNSRDFLRHFGCPVGSPMNPG |
| OR APPLICATION NUMBER: OR FILING DATE: 1998-05- OR PLING DATE: 1998-05- OR APPLICATION NUMBER: OR PLING DATE: 1998-05- OR APPLICATION NUMBER: | ry Match t Local Similar: ches 267; Cons | 2 EAETGSSV : 7 ELGAGSNVGF | 56 KPECII | 6 KELLE | 176 PEC | 233 EDYLD : 233 DYYL-1 | 292 TSEAM) : 292 DEEKIN | 352 RILGSERKKT : 346 ELINRTEPSI | 411 ESALPY : : 406 DDALG | 471 AVLAKV : : 465 AIYDMI | 531 FTNPT7 525 SMTPQ1 | 591 RKYDK : 584 REYDKI | 651 REAFR. : : 642 KAAYN | 711 SPPQFF : 699 SPARFF |
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TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic IITLE OF INVENTION: Acids Encoding the Same
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CURRENT APPLICATION NUMBER: US/09/978,192A
CURRENT FILING DATE: 2001-01-5
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PRIOR PILING DATE: 1997-11-03
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PRIOR FILING DATE: 1997-11-13
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RESULT 8
US-09-978-192A-526
Sequence 526, Application US/09978192A
Fatent No. US20020177553A1
GENERAL INFORMATION:
APPLICANT: Ashkenazi, Avi
APPLICANT: Baker Kevin P.
APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
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Godowski, Paul J.
Grimaldi, J. Christopher
Gurney, Austin L.
Hillan, Renneth J
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Paoni, Nicholas F.
Roy, Margaret Ann
Shelton, David L.
Stewart, Timothy A.
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Filvaroff, Ellen
Fong, Sherman
Gao, Wei-Qiang
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Gerritsen, Mary E.
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Kuo, Sophia S.
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Tumas, Daniel
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PRIOR APPLICATION NUMBER: 60/07963
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  176 P---EGVWSERKFSLLQTLATFRGQYSNSVFIRLYVSPDDKASNEHILKLDQATLSLAVR 232
                                                                                  233 EDYLDNSTEAKSYRDALYKFMVDTAVLLGANSSRAEHDMKSVLRLEIKIAEIMIPH-ENR 291
                                                                                                                                                            292 TSEAMYNKMINISELSAMIPQFDWLGYIKKVIDTRLYPHLKDISPSENVVVRVPQYFKDLF 351
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Grimaldi, J. Christopher
Gurney, Austin L.
Hillan, Kenneth J
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Shelton, David L.
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Filvaroff, Ellen
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Kuo, Sophia S.
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Botstein, David
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Gao, Wei-Qiang
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Matches 267; Conservative 145; Mismatches 307; Indels 40; Gaps
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APPLICATION NUMBER: 60/083545
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           TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic TITLE OF INVENTION: Acids Encoding the Same File Reference: P2630P1G5 CURRENT APPLICATION NUMBER: US/09/999,832A CURRENT FILING DATE: 2001-10-24 PRIOR APPLICATION NUMBER: 09/918585 PRIOR FILING DATE: 2001-07-30
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FILING DATE: 1998-03-25
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                                                                                                                                                                                        699 SPARFRVLGTLSNSRDFLRHFGCPVGSPMNPG-QLCEVW 736
                                                                                                                                              711 SPPQFRVNGAISNSEEFQXAFNCPPNSTMNRGMDSCRLW 749
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CURRENT APPLICATION NUMBER: US/09/978,189
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Grimaldi, J. Christopher
Gurney, Austin L.
Hillan, Kenneth J
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Roy, Margaret Ann
Shelton, David L.
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Filvaroff, Ellen
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Gao, Wei-Qiang
Gerber, Hanspeter
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Desnoyers, Luc
Eaton, Dan
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Ashkenazi, Avi APPLICANT: Baker Kevin P.
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Williams, P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  116 KELLEKSISRRRDTEAIQKAKILYSSCMNEKAIEKADAKPLHILRHSPFRWPVLESNIG 175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          127 KHLLENT-TFNSSSEAEOKTORFYLSCLOVERIEELGAOPLRDLI------EKIG 174
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      292 TSEAMYNNOMNISELSAMIPQFDWLGYIKKVIDTRLYPHLKDISPSENVVVRVPQYFKDLF 351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               292 DEEKIYHKMSISELQALAPSMDWLEFLSFLLSP-----LELSDSEPVVVYGMDYLQQVS 345
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APPLICATION NUMBER: 60/079728 FILING DATE: 1998-03-27 APPLICATION NUMBER: 60/079786 APPLICATION NUMBER: 60/079923 FILING DATE: 1998-03-30 APPLICATION NUMBER: 60/080105 APPLICATION NUMBER: 60/080165 FILING DATE: 1998-03-31 APPLICATION NUMBER: 60/080194 APPLICATION NUMBER: 60/081070 FILING DATE: 1998-04-08 FILING DATE: 1998-04-09 APPLICATION NUMBER: 60/081229 FILING DATE: 1998-04-15 APPLICATION NUMBER: 60/081819 FILING DATE: 1998-04-15 APPLICATION NUMBER: 60/082568 APPLICATION NUMBER: 60/082569 FILING DATE: 1998-04-21 APPLICATION NUMBER: 60/082704 FILING DATE: 1998-04-22 APPLICATION NUMBER: 60/082796 FILING DATE: 1998-03-27 APPLICATION NUMBER: 60/079663 FILING DATE: 1998-03-27 APPLICATION NUMBER: 60/079920 FILING DATE: 1998-03-30 ILING DATE: 1998-03-31 PPLICATION NUMBER: 60/080107 PPLICATION NUMBER: 60/080328 APPLICATION NUMBER: 60/080333 FILING DATE: 1998-04-01 APPLICATION NUMBER: 60/080334 APPLICATION NUMBER: 60/081049 FILING DATE: 1998-04-08 APPLICATION NUMBER: 60/081071 APPLICATION NUMBER: 60/081195 60/081955 APPLICATION NUMBER: 60/081817 LICATION NUMBER: 60/081952 APPLICATION NUMBER: 60/081838 FILING DATE: 1998-04-22 APPLICATION NUMBER: 60/082804 APPLICATION NUMBER: 60/082700 PPLICATION NUMBER: 60/082797 FILING DATE: 1998-03-25 APPLICATION NUMBER: 60/079656 FILING DATE: 1998-03-26 APPLICATION NUMBER: 60/080327 APPLICATION NUMBER: 60/081203 APPLICATION NUMBER: 60/078939 FILING DATE: 1998-03-20 APPLICATION NUMBER: 60/079664 FILING DATE: 1998-03-27 APPLICATION NUMBER: 60/079689 APPLICATION NUMBER: 60/079294 1998-04-08 1998-04-08 1998-04-09 1998-04-15 FILING DATE: 1998-04-23 FILING DATE: 1998-03-20 1998-03-31 LING DATE: 1998-04-01 1998-04-09 1998-04-15 1998-04-21 1998-04-22 1998-04-22 1998-04-01 1998-04-15 FILING DATE: 1998-03-27 8-60-8661 APPLICATION NUMBER: LING DATE: LING DATE: LING DATE: LING DATE: ILING DATE: REPRESENTATION OF PRESENTATION
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| PRICE APPLICATION NUMBER: (0/08332 | PRICE APPLICATION NUMBER: (0/08322 | PRICE APPLICATION NUMBER: (0/08322 | PRICE APPLICATION NUMBER: (0/08322 | PRICE APPLICATION NUMBER: (0/08322 | PRICE APPLICATION NUMBER: (0/08322 | PRICE APPLICATION NUMBER: (0/08326 | PRICE APPLICATION NUMBER: (0/08326 | PRICE APPLICATION NUMBER: (0/08326 | PRICE APPLICATION NUMBER: (0/08326 | PRICE APPLICATION NUMBER: (0/08326 | PRICE APPLICATION NUMBER: (0/08326 | PRICE APPLICATION NUMBER: (0/08326 | PRICE APPLICATION NUMBER: (0/08326 | PRICE APPLICATION NUMBER: (0/08326 | PRICE APPLICATION NUMBER: (0/08326 | PRICE APPLICATION NUMBER: (0/08326 | PRICE APPLICATION NUMBER: (0/08326 | PRICE APPLICATION NUMBER: (0/08326 | PRICE APPLICATION NUMBER: (0/08326 | PRICE APPLICATION NUMBER: (0/08326 | PRICE APPLICATION NUMBER: (0/08326 | PRICE APPLICATION NUMBER: (0/08326 | PRICE APPLICATION NUMBER: (0/08326 | PRICE APPLICATION NUMBER: (0/08326 | PRICE APPLICATION NUMBER: (0/08326 | PRICE APPLICATION NUMBER: (0/08326 | PRICE APPLICATION NUMBER: (0/08426 | PRICE APPLICATION NUMBER: (0/08436 | PRICE APPLICATION NUMBER: (0/08436 | PRICE APPLICATION NUMBER: (0/08436 | PRICE APPLICATION NUMBER: (0/08436 | PRICE APPLICATION NUMBER: (0/08436 | PRICE APPLICATION NUMBER: (0/08436 | PRICE APPLICATION NUMBER: (0/08436 | PRICE APPLICATION NUMBER: (0/08436 | PRICE APPLICATION NUMBER: (0/08436 | PRICE APPLICATION NUMBER: (0/08439 | PRICE APPLICATION NUMBER: (0/08439 | PRICE APPLICATION NUMBER: (0/08439 | PRICE APPLICATION NUMBER: (0/08439 | PRICE APPLICATION NUMBER: (0/08439 | PRICE APPLICATION NUMBER: (0/08439 | PRICE APPLICATION NUMBER: (0/08439 | PRICE APPLICATION NUMBER: (0/08439 | PRICE APPLICATION NUMBER: (0/08439 | PRICE APPLICATION NUMBER: (0/08439 | PRICE APPLICATION NUMBER: (0/08439 | PRICE APPLICATION NUMBER: (0/08439 | PRICE APPLICATION NUMBER: (0/08439 | PRICE APPLICATION NUMBER: (0/08439 | PRICE APPLICATION NUMBER: (0/08439 | PRICE APPLICATION NUMBER: (0/08439 | PRICE APPLICATION NUMBER: (0/08439 | PRICE APPLICATION NUMBER: (0/08439 | PRICE APP
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KELLEKSISRRRDTEAIQKAKILYSSCMNEKAIEKADAKPLLHILRHSPRWPVLESNIG 175
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                                                                                                                                                                                                                                       175 GWNIIGPWDQDNF--MEVLKAVAGIYRATPFFIVYISADSKSSNSNVIQVDQSGLFLPSR 232
                                                                                                                                                                                                                                                                                                                                                                                                                                               292 DEEKIYHKMSISELQALAPSMDWLEFLSFLLSP-----LELSDSEPVVVYGMDYLQQVS 345
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56 KPECIEAAAAILSKVNLSVDPCDNFFRFACDGWISNNPIPEDMPSYGVYPWLRHNVDLKL 115
                                                                                                                                                                                                    176 P---EGVWSERKFSLLQTLATFRGQYSNSVFIRLYVSPDDKASNEHILKLDQATLSLAVR 232
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Goddard, Audrey,
Goddowski, Paul J.
Godmaldi, J. Christopher
Gurney, Austin L.
Hillan, Kenneth J
Kljavin, Ivar J.
Kuo, Sophia S.
                  Sequence 526, Application US/09978608A
                                    US20030045462A1
                                                                                                                                                                                                                        Gerber, Hanspeter
Gerritsen, Mary E.
                                                                                                                                                     Ferrara, Napoleon
Filvaroff, Ellen
                                                                                                      Botstein, David
                                                                                                                     Desnoyers, Luc
Eaton, Dan
                                                                                                                                                                                    Fong, Sherman
Gao, Wei-Qiang
                                                                                    Baker Kevin P.
                                                                    APPLICANT: Ashkenazi, Avi
                                  Publication No. US20
GENERAL INFORMATION:
US-09-978-608A-526
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Query Match 30.9%; Score 1223; DB 11; Length 736;
Best Local Similarity 35.2%; Pred. No. 1.3e-105;
Matches 267; Conservative 145; Mismatches 307; Indels 40; Gaps 18;
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                                                                                                                     APPLICANT: Tumas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
                                                                                                                                                                                                                                                   FILE REFERENCE: P2630PIC22
CURRENT APPLICATION VUMBER: US/09/978,608A
CURRENT PLING DATE: 2001-10-16
NUMBER OF SEQ ID NOS: 624
Prior Application removed - See File Wrapper or Palm SEQ ID NO 526
LENGTH: 736
                                                                                     Shelton, David L.
Stewart, Timothy A.
               Pan, James,
Paoni, Nicholas F.
Roy, Margaret Ann
Napier, Mary A.
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US-09-978-608A-526
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233 DYYL-NRTANEKVLTAYLDYMEELGMLLGGRPTSTREQMQQVLELEIQLANITVPQDQRR 291
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Roy, Margaret Ann
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Filvaroff, Ellen
Fong, Sherman
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Gerber, Hanspeter
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Botstein, David
Desnoyers, Luc
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Napier, Mary A.
Pan, James;
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                            711 SPPQFRVNGAISNSEEFQKAFNCPPNSTMNRGMDSCRLW 749
                                                                  699 SPARFRVLGTLSNSRDFLRHFGCPVGSPMNPG-QLCEVW 736
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CURRENT FILING DATE: 2001-10-16
NUMBER OF SEQ ID NOS: 624
Prior Application removed - See File Wrapper or Palm
                                                                                                                                                          Sequence 526, Application US/09978585A Publication No. US20030049633A1
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Gurney, Austin L.
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Stewart, Timothy A.
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Filvaroff, Ellen
Fong, Sherman
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Botstein, David
Desnoyers, Luc
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Kuo, Sophia S.
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GENERAL INFORMATION:
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                                                                                                                                 352 RILGSERKKTIANYLVWRMVYSRIPNLSRRFQYRWLEFSRVIQGT-TTLLPQWDKCVNFI 410
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                                                              292 DEEXIYHKMSISELQALAPSMDWLEFLSFLLSP-----LELSDSEPVVYGMDYLQQVS 345
                                                                                                                                                                                                                                                                  411 ESALPYVVGKMFVDVYFQEDKKEMMEELVEGVRWAFIDMLEKENEWMDAGTKRKAKEKAR 470
                                                                                                                                                                                                                                                                                                                                 406 DDALGFALGSLFVXATFDROSKEIAEGMISEIRTAFEEAL-GOLVWMDEKTRQAAKEKAD 464
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OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic

OF INVENTION: Acids Encoding the Rama
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CURRENT APPLICATION NUMBER: US/09/978,191A
CURRENT FILING DATE: 2001-10-15
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; Publication No. US20030050239A1
; GENERAL INFORMATION:
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Gurney, Austin L.
Hillan, Kenneth J
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Williams, P. Mickey
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PRIOR FILING DATE: 2001-07-30

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PRIOR PELICATION UNDRESS: 60/66149

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30.9%; Score 1223; DB 11; Length 736;
Best Local Similarity 35.2%; Pred. No. 1.3e-105;
Matches 267; Conservative 145; Mismatches 307; Indels 40;
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PRIOR FILING DATE: 1998-05-13
PRIOR PAPPLICATION NUMBER: 60/085323
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PRIOR FILING DATE: 1998-05-15
PRIOR PAPLICATION NUMBER: 60/08582
PRIOR PILING DATE: 1998-05-15
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PRIOR APPLICATION NUMBER: 60/08559
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APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
TITLE OF INVENTION: Acids Encoding the Same
699 SPARFRVLGTLSNSRDFLRHFGCPVGSPMNPG-QLCEVW 736
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PRIOR APPLICATION NUMBER: 09/918565
PRIOR PILING DATE: 2001-07-30-9
PRIOR PILING DATE: 2001-07-30-9
PRIOR PILING DATE: 1997-10-17
PRIOR PELING DATE: 1997-10-17
PRIOR APPLICATION NUMBER: 60/064249
PRIOR FILING DATE: 1997-11-03
PRIOR PELING DATE: 1997-11-13
PRIOR APPLICATION NUMBER: 60/065364
PRIOR PELING DATE: 1997-11-21
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                                                                                                                                     Sequence 526, Application US/09978403A publication No. US20030050240Al GENERAL INFORMATION:
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Grimaldi, J. Christopher
Gurney, Austin L.
Hillan, Kenneth J
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PRIOR FILING DATE: 1998-03-20
PRIOR APPLICATION NUMBER: 60/078939
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FILING DATE: 1998-03-25
APPLICATION NUMBER: 60/079656
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Shelton, David L.
Stewart, Timothy A.
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Paoni, Nicholas F.
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Filvaroff, Ellen
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Kuo, Sophia S.
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APPLICANT: Baker Kevin P.
APPLICANT: Botstein, David
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56 KPECIEAAAAILSKVNLSVDPCDNFFRFACDGWISNNPIPEDMPSYGVYPWLRHNVDLKL 115
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R APPLICATION NUMBER: 60/084600

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R APPLICATION NUMBER: 60/084643
  APPLICATION NUMBER: 60/083496 FILING DATE: 1998-04-29 FILING DATE: 1998-04-20
                                                                                     APPLICATION NUMBER: 60/083499
FILING DATE: 1998-04-29
APPLICATION NUMBER: 60/083545
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APPLICATION WIMBER: 60/085338
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APPLICATION NUMBER: 60/085323
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PRIOR APPLICATION WUMBER: 60/085704
PRIOR FLING DATE: 1998-05-15
PRIOR APPLICATION NUMBER: 60/085697
APPLICATION NUMBER: 60/083495
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APPLICATION NUMBER: 60/083554
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APPLICATION NUMBER: 60/083500
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FILING DATE: 1998-04-30
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FILING DATE: 1998-05-05
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FILING DATE: 1998-05-06
APPLICATION NUMBER: 60/08441
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APPLICATION NUMBER: 60/084640
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APPLICATION NUMBER: 60/084598
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APPLICATION NUMBER: 60/085339
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APPLICATION NUMBER: 60/085582
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APPLICATION NUMBER: 60/085700
FILING DATE: 1998-05-15
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APPLICATION UNMBER: 60/085580
FILING DATE: 1998-05-15
APPLICATION NUMBER: 60/085573
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176 P---EGVWSERKFSLLQTLATFRGQYSNSVFIRLYVSPDDKASNEHILKLDQATLSLAVR 232
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127 KHLLENT-TFNSSSEAEQKTQRPYLSCLQVERIEELGAQPLRDLI------EKIG 174
                                                                                                            175 GWNITGPWDQDNF--MEVLKAVAGTYRATPFFTVYISADSKSSNSNVIQVDQSGLFLPSR 232
                                                                                                                                                                                                                                         292 TSEAMYNKWANISELSAMIPQFDWLGYIKKVIDTRLYPHLKDISPSENVVVRVPQYFKDLP 351
                                                                                                                                                      233 EDYLDNSTEAKSYRDALYKFMVDTAVLLGANSSRAEHDMKSVLRLEIKIAEIMIPH-ENR 291
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Grimaldi, J. Christopher
Gurney, Austin L.
Hillan, Kenneth J
Kljavin, Ivar J.
Kuo, Sophia S.
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Shelton, David L.
Stewart, Timothy A.
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Gerritsen, Mary E.
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Filvaroff, Ellen
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APPLICANT: Baker, Kevin P.
APPLICANT: Bustein, David
APPLICANT: Desnoyers, Luc
APPLICANT: Baton, Dan
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TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic TITLE OF INVENTION: Acids Encoding the Same File REFERENCE: P2630991C25
CURRENT APPLICATION NUMBER: US/09/978,564A
CURRENT FILING DATE: 2001-10-16
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                               Williams, P. Mickey Wood, William I.
Tumas, Daniel
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| 1998-04-08 NUMBER: 60/081049 1998-04-08 1998-04-08 1998-04-08 1998-04-08 NUMBER: 60/081203 1998-04-09 1998-04-09 1998-04-09 1998-04-09 1998-04-09 1998-04-09 | 98-04-15 98-04-15 98-04-15 98-04-15 98-04-15 98-04-15 98-04-15 98-04-15 98-04-15 98-04-15 98-04-15 98-04-25 98-04-25 98-04-25 98-04-25 98-04-25 98-04-25 98-04-25 98-04-25 98-04-25 98-04-25 98-04-25 98-04-25 98-04-25 98-04-25 | 28 - 0 - 0 - 0 - 0 - 0 - 0 - 0 - 0 - 0 - |
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R APPLICATION NUMBER: 60/085339
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APPLICATION NUMBER: 60/084627
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Sequence 16877, A
Sequence 15, Appl
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Sequence 30164, A
Sequence 573, App
Sequence 26336, A
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Sequence 3801, Ap
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                                                                               December 3, 2003, 07:21:37; Search time 0.627451 Seconds (without alignments) 1078.927 Million cell updates/sec
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/cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PeTUS_COMB.pep:*
/cgn2_6/ptodata/1/iaa/PeTUS_COMB.pep:*
GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-107-552A-381

US-09-107-552A-381

US-09-252-991A-301626

US-09-252-991A-20184

US-09-252-991A-20184

US-09-252-991A-20184

US-09-252-991A-18492

US-09-252-991A-18492

US-09-252-991A-18710

US-09-252-991A-18710

US-09-252-991A-18710

US-09-252-991A-18710

US-09-252-991A-17641

US-09-252-991A-17641

US-09-252-991A-17641

US-09-252-991A-17641

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US-09-252-991A-17641

US-09-252-991A-17641

US-09-252-991A-17641

US-09-252-991A-17641

US-09-31899-9

US-09-251-891-9

US-09-271-891-9

US-09-271-891-9

US-09-271-891-9

US-09-271-891-9

US-09-271-891-9

US-09-271-891-9
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US-09-602-877A-99
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                                                                                                                                                                                                                                               328717 segs, 42310858 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SUMMARIES
                                                         OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                Listing first 45 summaries
                                                                                                                                                                                                                     Gapop 10.0 , Gapext 0.5
                                                                                                                                                                         1 VLTVIAQQTTSQGLLS 16
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                                                                                                                                                                                                                                                                                                                                                                  Maximum Match 100%
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Maximum DB seq length: 2000000000
                                                                                                                                                                                                                                                                                                                                                    Post-processing: Minimum Match 0%
                                                                                                                                       US-09-913-955A-4
72
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Match Length DB
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749
2004
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2004
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1034
1627
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453
459
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0.0000884774
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45.1
45.1
45.1
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Perfect score:
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                                                                                                                                                                         Sequence:
                                                                                                                                                                                                                                               Searched:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    lesult
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Sequence 424, Application US/09198452A

Patent No. 6559294

GENERAL INFORMATION:

APPLICANT: Griffals, R.

TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnosis, previsitle REPERENCE: 9710-003-999

CURRENT APPLICATION NUMBER: 105/198,452A

CURRENT FILING DATE: 1998-11-24

NUMBER OF SEQ ID NOS: 6849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Lynn A Doucette-Stamm and David Bush
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
ENTEROCOCCUS FAECIUM FOR DIAGNOSTICS AND THERAPEUTICS
   Sequence 28029, A Sequence 16, Appl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .
                                                                      sequence 5629. Appleaguence 17419, A Sequence 3301, A Sequence 2, Appli Sequence 6805, Appli Sequence 6805, Appli Sequence 6805, Appli
                                                                                                                                                                                                                                         5839, Ap
28610. A
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28398, A
                                                                                                                                                                                Sequence 6253, Ap
Sequence 20913, A
 Sequence 403,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: GENOME THERAPEUTICS CORPORATION STREET: 100 Beaver Street
                           US-09-252-991A-28029
US-08-685-99216
US-09-14-955-16
US-09-328-352-5629
US-09-328-352-5629
US-09-134-001C-301
US-08-812-586-2
US-09-535-882A-2
US-09-535-882A-2
US-09-535-882A-2
US-09-252-991A-2013
US-09-252-991A-2013
US-09-252-991A-2013
US-09-252-991A-2013
US-09-252-991A-2013
US-09-252-991A-2013
US-09-252-991A-2013
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                                                                                                                                                                                                                                                                                                    ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: CD/ROM ISO9660
COMPUTER: FOC
OPERATING SYSTEM: <Unknown>
SOFTWARE: ASCII
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  , Sequence 3801, Application US/09107532A
; Patent No. 6583275
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT ORGANISM: Chlamydia pneumoniae
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 7310
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         162 LLSLVSOMTTSQG 174
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2006
2006
2007
2007
3007
3107
3107
3107
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us-09-913-955a-4.rai

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; Sequence 30164, Application US/09252991A; Patent No. 6551795; GENERAL INFORMATION:
                                                                                                                                                                               Sequence 30426, Application US/09252991A Patent No. 6551795
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US-09-198 452A-573
Sequence 573, Application US/09198452A
Patent No. 6552294
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Pseudomonas aeruginosa
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 VLTVIAQQTTSQGLL 15
                                                                  1565 TELVÖKSTSEGLL 1577
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 33142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 55...
Best Local 8, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 33142
                          3 TVIAQQTTSOGLL 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity 58.3
Matches 7; Conservative
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                                                                                                                                                          US-09-252-991A-30426
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US-09-252-991A-30164
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LENGTH: 361
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LENGTH: 454
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OTHER INFORMATION: Description of Artificial Sequence: Full length
OTHER INFORMATION: utrophin construct; Xaa = unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
51.4%; Score 37; DB 4; Length 3433;
Best Local Similarity 53.8%; Pred. No. 7.8e+02;
Matches 7; Conservative 4; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Match 51.4%; Score 37; DB 4; Length 206; Local Similarity 61.5%; Pred. No. 25; ess 8; Conservative 1; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Tinsley, Jonathon M
APPLICANT: Davies, Kay E
TITLE OF INVENTION: Utrophin gene expression
FILE REFERENCE: 620-42
CURRENT APPLICATION NUMBER: US/09/091,5018
CURRENT FILING DATE: 1998-06-18
PRIOR APPLICATION NUMBER: PCT/GB96/03156
PRIOR FILING DATE: 1996-12-19
PRIOR FILING DATE: 1995-12-19
PRIOR FILING DATE: 1996-12-19
PRIOR APPLICATION NUMBER: GB 9525962,8
PRIOR FILING DATE: 1996-07-26
PRIOR FILING DATE: 1996-07-26
NUMBER OF SEQ ID NOS: 15
SOFTWARE: PATENTIN NUMBER: GB 9622174.2
                                                                                                                                                                        NAME: Ariniello, Pamela Deneke
REGISTRATION NUMBER: 40.489
REFERENCE/DOCKET NUMBER: GTC-012
TELECOMMUNICATION INFORMATION:
TELEPHONE: (781)893-5007
TELEFAX: (781)893-5007
INFORMATION FOR SEQ ID NO: 3801:
SEQUENCE CHRACATERISTICS:
LENGTH: 206 amino acids
TYPE: amino acid
APPLICATION NUMBER: US/09/107,532A FILING DATE: 30-Jun-1998 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LOCATION: (B) LÖCATION 1...206
SEQUENCE DESCRIPTION: SEQ ID NO: 3801:
                                                             APPLICATION NUMBER: 60/085,598
FILING DATE: 14 May 1998
APPLICATION NUMBER: 60/051571
FILING DATE: July 2, 1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: YES
ORIGHAL SOURCE:
ORGANISM: Enterococus faecium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-091-501B-10; Sequence 10, Application US/09091501B Sequence 10, Application US/09091501B; General No. 6518413; General INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        misc feature (B) LOCATION 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY:
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GENERAL INFORMATION:

APPLICANT: Marc J. Rubenfield et al.

APPLICANT: Marc J. Rubenfield et al.

APPLICANT: Marc J. Rubenfield et al.

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS

TITLE OF INVENTION: ABERDOINOSA FOR DIAGNOSTICS AND THERAPEUTICS

FILE REFERENCE: 107196.136

CURRENT FILING DATE: 1999-02-18

PRIOR PLILING DATE: 1999-02-18

PRIOR PLILING DATE: 1999-02-18

PRIOR PLILING DATE: 1999-02-18

PRIOR FILING DATE: 1999-07-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATE MALC J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AUGUSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
PRIOR APPLICATION NUMBER: US/09/074,788
PRIOR APPLICATION NUMBER: US/09/07-180
PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1999-07-27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 50.0%; Score 36; DB 4; Length 454; 58.3%; Pred. No. 1e+02; tive 2; Mismatches 3; Indels
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Best Local Similarity 43.8
Matches 7; Conservative
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Matches 6; Conserv
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APPLICANT: Griffais, R.

TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence and polypeptides, fragments
TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnosis, preve
TITLE OF INVENTION: and treatment of infection
FILE REFERENCE: 9710-003-999
CURRENT APPLICATION NUMBER: US/09/198,452A
CURRENT FILING DATE: 1998-11-24
NUMBER OF SEQ ID NOS: 6849
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Section No. 6562958

GRNERAL INFORMATION:
GRNERAL INFORMATION:
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
TITLE OF INVENTION: BAUMANNI FOR DIAGNOSTICS AND THERAPEUTICS
TITLE OF INVENTION: BAUMANNI FOR DIAGNOSTICS AND THERAPEUTICS
CURRENT APPLICATION NUMBER: US/09/328,352
CURRENT FILING DATE: 1999-06-04
NUMBER OF SEQ ID NOS: 8252
SEQ ID NO 4457
LENGTH: 705
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 26336, Application US/09252991A
Patent No. 6551795
Patent No. 6551795
Patent No. 6551795
TITLE OF INVEXTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVEXTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVEXTION: NUCLEIC ACID AND THERAPEUTICS
CURRENT APPLICATION NUMBER: US/09/252,991A
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1999-07-27
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                                                                                                                                                                                                                                                                                                                                                                                    3; Mismatches
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                                                                                                                                                                                                                                          TYPE: PRT
CRGANISM: Chlamydia pneumoniae
US-09-198-452A-573
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Best Local Similarity 53....
Best Local Similarity 53....
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Best Local Similarity 50.0%
Matches 8; Conservative
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JS-09-328-352-4457
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LENGTH: 625
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                                                                                                                                                                                            SEQ ID NO 573
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Sequence 19492, Application US/09252991A
Sequence 19492, Application US/09252991A
Sequence 19492, Application US/09252991A
Sequence 19492, Application US/09252991A
Sequence 19492, Application US/09252991A
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
CURRENT APPLICATION NUMBER: US/09/252,991A
FRIOR FILING DATE: 1998-02-18
PRIOR FILING DATE: 1998-02-18
PRIOR FILING DATE: 1998-02-18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Clean, Matthew
APPLICANT: Lubbers, Mark W
APPLICANT: Lubbers, Mark W
APPLICANT: Debkers, Jamer W
APPLICANT: Debkers, Jamer W
TITLE OF INVENTION: Polynucleotides, materials incorporating
TITLE OF INVENTION: Polynucleotides, materials incorporating
TITLE OF INVENTION: Polynucleotides, materials incorporating
TITLE REFERENCE: 1048011
CURRENT APPLICATION NUMBER: US/09/724,623
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 124
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           48.6%; Score 35; DB 4; Length 803; 46.2%; Pred. No. 3.1e+02; tive 5; Mismatches 2; Indels
43.8%; Pred. No. 2.7e+02;
tive 4; Mismatches 5; Indels
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US-09-724-623-106
Sequence 106, Application US/09724623
Patent No. 6476209
GENERAL INFORMATION:
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20 TVLSQLTTAQKNAGLLA 36
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                                                                                            1 VLTVIAQOTTSQGLLS
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Matches 9; Conservative
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Sequence 17641, Application US/09252991A
Patent No. 6551795
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
TITLE OF INVENTION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR PAPLICATION NUMBER: US 60/074,788
PRIOR PELICATION NUMBER: US 60/094,190
PRIOR PELICATION NUMBER: US 60/094,190
PRIOR PELICATION NUMBER: US 60/094,190
PRIOR PELICATION NUMBER: US 60/094,190
PRIOR PILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 1761
                                                                                                                           0; Gaps
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                                                                 45.8%; Score 33; DB 4; Length 438; 46.7%; Pred. No. 3.5e+02; ative 3; Mismatches 5; Indels
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Patent No. 5925566
ENDEAL INCAMATION:
APPLICANT: Davis, Roger J.
APPLICANT: Galchew-Gargova, Zoya
TITLE OF INVENTION: NON-ACTIVATED RECEPTOR COMPLEX
TITLE OF INVENTION: PROTEINS AND USES THEREOF
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
45.8%; Score 33; DB 4; I
Best Local Similarity 46.2%; Pred. No. 3.7e+02;
Matches 6; Conservative 3; Mismatches 4;
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FSELSEQ for Windows Version 2.0
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APPLICATION WUMBER: US/08/870,518
FILING DATE: 06-UNA-1997
PRIOR APPLICATION DATA:
APPLICATION WUMBER: 60/019,219
FILING DATE: 06-UNA-1996
ATTORNEY/AGENT INFORMATION:
ORGANISM: Pseudomonas aeruginosa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Pseudomonas aeruginosa
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                                                                                                 Best Local Similarity 46.7%
Matches 7; Conservative
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STREET: ZZZ
TTY; Boston
; OKGANISH: ...
US-09-252-991A-18710
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US-08-870-518-4
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                                                                             Query Match
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APPLICANT: Gary L. Breton et al.
TITLE OF INVENTION: NUCLEIC ACID AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
TITLE OF INVENTION: BAUNANNI FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: GTC99-03PA
CURRENT APPLICATION NUMBER: US/09/328,352
CURRENT FILING DATE: 1999-06-04
NUMBER OF SEQ ID NOS: 9252
              APPLICANT: MAIC J. Rubenfield et al.

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS FILE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS FILE REPERBNCE: 107196.136

CURRENT APPLICATION NUMBER: US/09/252,991A

CURRENT FILING DATE: 1999-02-18

PRIOR FILING DATE: 1999-02-18

PRIOR FILING DATE: 1998-02-18

PRIOR FILING DATE: 1998-07-27

NUMBER OF SEQ ID NOS: 33142

SEQ ID NO 28921
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APPLICANT: MAC J. Rubenfield et al.

TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS

TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS

TITLE OF INVENTION: ACRUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS

CURRENT APPLICATION NUMBER: US/09/252,991A

CURRENT APPLICATION NUMBER: US/09/252,991A

PRIOR FILING DATE: 1998-02-18

PRIOR FILING DATE: 1998-02-18

PRIOR APPLICATION NUMBER: US 60/094,190

PRIOR APPLICATION NUMBER: US 60/094,190

NUMBER OF SEQ ID NOS: 33142
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47.2%; Score 34; DB 4; Length 1034;
Best Local Similarity 54.5%; Pred. No. 6.5e+02;
Matches 6; Conservative 3; Mismatches 2; Indels
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US-09-252-991A-18710
; Sequence 18710, Application US/09252991A
; Patent No. 6551795
; GENERAL INFORMATION:
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8-09-328-352-6604
; Sequence 6604, Application US/09328352
; Patent No. 6562958
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US-09-328-352-6604
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GENERAL INFORMATION:
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US-09-252-991A-28921
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LENGTH: 438
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| REGISTRATION NUMBER: 32,983 | REGISTRATION NUMBER: 04020/102001 | TELECOMMUNICATION INFORMATION: TELECOMMUNICATION INFORMATION: TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX: 200154 | TELEPAX:
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Sequence 340, App
Sequence 10486, App
Sequence 10486, A
Sequence 2, Appli
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Sequence 5842, Ap
Sequence 12564, A
Sequence 11, Appl
Sequence 477, Appl
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Sequence 27413, A
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Sequence 1641, Ap
Sequence 12525, A
                                                                                                                                                                  December 3, 2003, 07:21:37; Search time 10.6248 Seconds (without alignments) 280.074 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Published Applications Asia (1902)

(gn2_6/ptodata/1/pubpaa/US07_PUBCQMB.pep.* (gn2_6/ptodata/1/pubpaa/US06_PUBG_PUB.pep.* (gn2_6/ptodata/1/pubpaa/US06_PUBG_PUB.pep.* (gn2_6/ptodata/1/pubpaa/US06_PUBGOMB.pep.* (gn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep.* (gn2_6/ptodata/1/pubpaa/US08_PUBGOMB.pep.* (gn2_6/ptodata/1/pubpaa/US08_PUBGOMB.pep.* (gn2_6/ptodata/1/pubpaa/US08_PUBGOMB.pep.* (gn2_6/ptodata/1/pubpaa/US08_PUBGOMB.pep.* (gn2_6/ptodata/1/pubpaa/US09_PUBGOMB.pep.* (gn2_6/ptodata/1/pubpaa/US09_PUBGOMB.pep.* (gn2_6/ptodata/1/pubpaa/US09_PUBGOMB.pep.* (gn2_6/ptodata/1/pubpaa/US09_PUBGOMB.pep.* (gn2_6/ptodata/1/pubpaa/US09_PUBGOMB.pep.* (gn2_6/ptodata/1/pubpaa/US09_PUBGOMB.pep.* (gn2_6/ptodata/1/pubpaa/US09_PUBGOMB.pep.* (gn2_6/ptodata/1/pubpaa/US00_PUBGOMB.pep.* (gn2_6/ptodata/1/pubpaa/US00_PUBGOMB.pep.* (gn2_6/ptodata/1/pubpaa/US00_NEW_PUB.pep.* (gn2_6/ptodata/1/pubpaa/US00_PUBCOMB.pep.* (gn2_6/ptodata/1/pubpaa/US00_P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           684280
GenCore version 5.1.6
Copyright (c) 1993 - 2003 Compugen Ltd.
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2 US-10-029-886-27413
0 US-09-816-247-84340
US-09-815-242-10486
4 US-10-047-539-2
5 US-10-047-539-2
6 US-10-310-002-3
6 US-10-310-002-3
7 US-10-310-002-47
5 US-10-310-002-47
5 US-10-310-002-47
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6 US-09-925-300-1661
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7 US-09-925-300-1661
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             684280 seqs, 185983659 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Published Applications AA:*
                                                                                                                OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                            BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                1 VLTVIAQQTTSQGLLS 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Maximum Match 100%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Minimum DB seq length: 0 Maximum DB seq length: 2000000000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Post-processing: Minimum Match 08
                                                                                                                                                                                                                                                                               US-09-913-955A-4
72
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Match Length DB
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45.8
8.8
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Searched:
                                                                                                                                                                           Run on:
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| 16 | 33 | īŲ. | v | 12 | -10-289-757-9 | 98, |
|---------|---------------|------|-----|----|-------------------|--------------|
| 17 | 33 | 'n | m | 10 | -09-738-626- | 429 |
| 18 | 33 | 45.8 | 476 | 10 | 'n | S |
| 19 | 33 | 'n | 5 | 10 | -09-813-398- | 26, |
| 20 | 33 | 'n | N | 10 | -09-738-626-4 | 4919 |
| 21 | 33 | 'n. | 0 | 12 | -10-199-869-6 | 6, 1 |
| 22 | 33 | 'n | S | 12 | -10-120-801-4 | 47, |
| 23 | 33 | ď. | 90 | 12 | -10-029-386- | 324(|
| 24 | 33 | 'n. | 00 | 13 | -10-331-061- | 7 '6 |
| 25 | 33 | ď. | S | 12 | 39-83 | 6 4 |
| 26 | ď. | S. | 07 | 10 | 09-801-368-11 | 110, |
| 27 | 32.5 | ď. | 32 | 10 | 39-801-368-1 | 114, |
| 28 | α. | 'n. | 53 | 70 | 39-801-368-10 | 104, |
| 29 | 32 | | 57 | 12 | 10-029-386-314 | 3145 |
| 30 | 32 | 44.4 | 4 | 15 | 10-156-761-97 | 9716 |
| 31 | 32 | 4, | 8 | 9 | 10-080-170-568 | 566 |
| 32 | 32 | 44.4 | æ | | 9-768-826-3 | equence 30, |
| 33 | 32 | 4 | 9 | | 9-745-288-9 | 66 |
| 34 | 32 | 44.4 | o, | S | 10-177-293-5 | Sequence 55, |
| 35 | 32 | 44.4 | 0 | | 9-989-722-40 | equence 403, |
| 36 | 32 | 4 | 206 | | -723-4 | e 403, |
| 3.7 | 32 | 4 | Ó | | 9-989-279-40 | equence 403, |
| 38 | 32 | 4 | Ö | | 9-989-727-40 | equence 403, |
| 39 | 32 | 4 | Ó | 10 | 5-09-989-731-40 | e 403 |
| 40 | 32 | 44.4 | ō | 10 | 5-09-989-732-40 | equence 403 |
| 41 | 32 | 4 | ō | 10 | S-09-991-073-40 | |
| 42 | 32 | 4 | ō | 10 | 09-990-442-40 | equence 403 |
| 43 | 32 | 4, | Ö | 10 | 09-991-163-40 | equence 403 |
| 44 | 32 | | ō | 10 | US-09-993-604-403 | equence 403 |
| 45 | 32 | | ō | 10 | 09-990-456-40 | e 4 |
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US-10-156-761-11650

US-10-156-761-11650

Sequence 11650, Application US/10156761

Publication No. US20030119018A1

GENERAL INFORMATION:

APPLICANT: IKEDA, HARUO

APPLICANT: ISHIKAMA, UNO

APPLICANT: SATAKAM, UNO

APPLICANT: SATAKAM, UNO

APPLICANT: SATAKAM, UNO

APPLICANT: SATAKAM, UNO

APPLICANT: SATAKAM, UNO

APPLICANT: SATAKAM, UNO

APPLICANT: SATAKAM, UNO

APPLICANT: SATAKAM, UNO

APPLICANT: MASHIRA

APPLICANT: MASHIRA

APPLICANT: MASHIRA

FILE CF INVENTION: NUMBER: US/10/156,761

CURRENT APPLICATION NUMBER: US/20-25-9

CURRENT APPLICATION NUMBER: US/20-25-9

PRIOR FILING DATE: 2001-05-30

PRIOR FILING DATE: 2001-05-30

NUMBER OF SEQ ID NOS: 15109

SEQ ID NO 11650

LENOTH: 453

TYPE: PRT

TYPE: PRT

CHENOTH: 453

ATYPE: PRT

CORSANISM: Streptomyces avermitilis

US-10-156-761-11650

QUETY MATCH

SO.0*; Pred: No. 3:10+02;

US-10-156-761-11650

QUETY MATCH

ATILALVRQUITSGEL 61

DD

47 ILALVRQUITSGEL 61
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0

RESULT 2 US-10-029-386-27413

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RESULT 5
US-09-815-242-10486
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                                                     US-09-815-242-4880
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Al-Garawi, Amal
APPLICANT: Miller, Charles
APPLICANT: Tomb, Jean-Francois
APPLICANT: Tomb, Jean-Francois
TITLE OF INVENTION: Identification of Polynucleotides
TITLE OF INVENTION: Encoding No. US20020115078Alel Helicobacter Polypeptides in the
                                                                             APPLICANT: Penn, Sharron G.
APPLICANT: Rank, David R.
APPLICANT: Rank, David K.
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR (TITLE OF INVENTION: EXPRESSION ANALYSIS TWO TITLE OF INVENTION: EXPRESSION ANALYSIS TWO CURRENT APPLICATION NUMBER: US/10/029,386
CURRENT APPLICATION NUMBER: 2011-2-20
CURRENT FILLE DEPENDENCE: 2001-12-20
SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      48.6%; Score 35; DB 12; Length 77;
40.0%; Pred. No. 60;
ative 5; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: MAP TO CHR22_105.0

OTHER INFORMATION: EXPRESSED IN HELD, SIGNAL = 1.1

OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 1.9

OTHER INFORMATION: EXPRESSED IN LING, SIGNAL = 1.9

OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 1.8

OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 1.9

OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 2.1

OTHER INFORMATION: EXPRESSED IN BARIN, SIGNAL = 2.1

OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 2.1

OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 2.1

OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 2.2

OTHER INFORMATION: SWISSPROT HIT: Q9VHB6, EVALUE 9.0000+00
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CURRENT APPLICATION NUMBER: US/09/881,752A
CURRENT FILING DATE: 2001-06-15
PRIOR APPLICATION NUMBER: US 08/833,457
PRIOR FILING DATE: 1997-04-01
NUMBER OF SEQ ID NOS: 370
SOFTWARE: FASTSEQ for Windows Version 4.0
Sequence 27413, Application US/10029386
Publication No. US20030194704A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 340, Application US/09881752A Patent No. US20020115078A1 PATENL INFORMATION:
APPLICANT: Kleanthous, Harold
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               173 ILTPIVAÇNVAÇGLES 188
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Best Local Similarity 40.0.
Best Local 6; Conservative
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Best Local Similarity 50.0%
Matches 8; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-10-029-386-27413
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                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO 27413
LENGTH: 77
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LENGTH: 236
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0; Gaps
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                                                                                                        APPLICANT: Oblean, Kari L.
APPLICANT: Zyskind, Judith W.
APPLICANT: Wall, Daniel
APPLICANT: Wall, Daniel
APPLICANT: Tawakck, John D.
APPLICANT: Yamamoro, Robert T.
APPLICANT: Yamamoro, Robert T.
APPLICANT: Yamamoro, Robert T.
APPLICANT: W. H. Howard
TITLE OF INVENTION: Identification of Essential Genes in
TITLE OF INVENTION: Prokaryotes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Wall, Daniel
APPLICANT: Travick, John D.
APPLICANT: Travick, John D.
APPLICANT: Yamamoto, Robert T.
APPLICANT: Yu. H. Howard
TITLE OF INVENTION: Identification of Essential Genes in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           48.6%; Score 35; DB 9; L
43.8%; Pred. No. 2.4e+02;
Live 5; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: ELITRA.011A
CURRENT APPLICATION NUMBER: US/09/815,242
CURRENT FILING DATE: 2001-03-21
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SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILE REFERENCE: BLITRA.011A
CURRENT APPLICATION NUMBER: US/09/815,242
CURRENT FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                          CUCKRAN FILING DATE: 2001-03-21
PRIOR FILING DATE: 2000-03-21
PRIOR FILING DATE: 2000-03-21
PRIOR PELING DATE: 2000-03-21
PRIOR PELING DATE: 2000-05-23
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/207,727
PRIOR APPLICATION NUMBER: 60/24,578
PRIOR PELING DATE: 2000-10-23
PRIOR PELING DATE: 2000-11-27
PRIOR PELING DATE: 2000-11-27
PRIOR APPLICATION NUMBER: 60/253,625
PRIOR APPLICATION NUMBER: 60/253,625
PRIOR APPLICATION NUMBER: 60/257,931
PRIOR APPLICATION NUMBER: 60/257,931
PRIOR PELING DATE: 2000-12-22
PRIOR FILING DATE: 2001-02-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Sequence 10486, Application US/09815242
; Patent No. USZ002006156941
; PENERAL INFORMATION:
; APPLICANT: Haselbeck, Robert
; APPLICANT: Ohlsen, Kari L.
; Sequence 4880, Application US/09815242; Patent No. US20020061569A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 60/191,078
PRIOR FILING DATE: 2000-03-21
PRIOR PELING DATE: 2000-05-23
PRIOR PELING DATE: 2000-05-23
PRIOR APPLICATION NUMBER: 60/207,727
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 VLTVIAQQTTSQGLLS 16
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Zyskind, Judith W.
                                                                                         APPLICANT: Haselbeck, Robert
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Best Local Similarity 45...
Best Local 7; Conservative
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LENGIH:
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Publication No. US20030138822A1
GENERAL INFORMATION:
APPLICANT: Glenn, Matthew
APPLICANT: Glenn, Mark W
APPLICANT: Glenn, Mark W
APPLICANT: Dekker, James
TITLE OF INVENTION: Polynucleotides and polypeptides, materials incorporating
TITLE OF INVENTION: Lhem, and methods for using them.
FILE REPERENCE: 11000.1048 ulc.1
CURRENT APPLICATION NUMBER: US/10/288,930
CURRENT FILING DATE: 2002-11-05
PRIOR FILING DATE: 2000-11-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: MOLLING, KARIN
APPLICANT: PAVLOVIC, JOVAN
APPLICANT: NAWRATH, MICHAEL
TITLE OF INVENTION: PHARMACEUTICAL COMPOSITIONS FOR TREATING OR PREVENTING
TITLE OF INVENTION: CANCER
FILE REFERENCE: VOS-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Gaps
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48.6%; Score 35; DB 14; Length 626;
Best Local Similarity 54.5%; Pred. No. 6.8e+02;
Matches 6; Conservative 3; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 258;
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48.6%; Score 35; DB 9; L
Best Local Similarity 43.8%; Pred. No. 2.4e+02;
Matches 7; Conservative 5; Mismatches 4;
PRIOR FILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: 60/242,578
PRIOR FILING DATE: 2000-10-23
PRIOR FILING DATE: 2000-110-23
PRIOR FILING DATE: 2000-111-27
PRIOR FILING DATE: 2000-11-27
PRIOR FILING DATE: 2000-11-27
PRIOR FILING DATE: 2000-12-22
PRIOR FILING DATE: 2000-12-22
PRIOR FILING DATE: 2010-12-26
NUMBER OF SEQ ID NOS: 14110
SOFTWARE: FastSEQ for Windows Version 4.0
SACTANARE: FastSEQ for Windows Version 4.0
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CURRENT FILING DATE: 2002-01-15
PRIOR APPLICATION NUMBER: EP 01 10 0914.9
PRIOR FILING DATE: 2001-01-16
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver: 2.1
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                                                                                                                                                                                                                                                                                                                                                                             ; ORGANISM: Enterococcus faecalis
US-09-815-242-10486
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 VLTVIAQQTTSQGLLS 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  80 VLNSLSELPTSQGILA 95
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US-10-047-539-2
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LENGTH: 626
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US-10-047-539-2
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APPLICANT: Emory University
TITLE OF INVENTION: Insulin-Responsive Sequence DNA Binding Protein-1 and Methods to
TITLE OF INVENTION: Regulate Insulin-Responsive Genes
TITLE OF INVENTION: Regulate Insulin-Responsive Genes
FILE REFRENCE: E056 1010
CURRENT APPLICATION WINBER: US/10/310,002
CURRENT FILING DATE: 2002-12-04
NUMBER OF SEQ ID NOS: 63
SOFTWARE: Patentin version 3.2
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APPLICANT: MAZOGUCHI, HIROSHI
APPLICANT: MADO, SELKO
APPLICANT: HAYASHI, MIKIRO
APPLICANT: HAYASHI, MIKIRO
APPLICANT: CCHIAI, KEIKO
APPLICANT: COKIAI, HARUHIKO
APPLICANT: TATEISHI, NAOKO
APPLICANT: TATEISHI, NASATO
APPLICANT: TATEISHI, NASATO
APPLICANT: TATEISHI, NASATO
APPLICANT: SENOH, AKINIRO
APPLICANT: TATEIA, MASATO
APPLICANT: COZAKI, AKIO
TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES
FILE REPERBUCE: 249-125
CURRENT APPLICATION NUMBER: US/09/738,626
CURRENT APPLICATION NUMBER: JP 99/377484
PRIOR FILING DATE: 2000-12-16
PRIOR APPLICATION NUMBER: JP 00/159162
PRIOR APPLICATION NUMBER: JP 00/159162
PRIOR APPLICATION NUMBER: JP 00/159162
PRIOR PRIUNG DATE: 2000-04-07
PRIOR PLING DATE: 2000-04-07
PRIOR PLING DATE: 2000-08-03
NUMBER OF SEQ ID NOS: 7059
PRIOR APPLICATION NUMBER: 60/148,801
PRIOR FLING DAFE: 1999-12-02
NUMBER OF SEQ ID NOS: 124
SOFTWARE: PASESEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-10-310-002-3
; Sequence 3, Application US/10310002
; Publication No. US20030125296A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                         ORGANISM: Lactobacillus rhamnosus
                                                                                                                                                                                                                                                                                                                                                                                                                                                  3 TVIAQQTTSQ---GLLS 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20 TVLSQLTTAQKNAGLLA 36
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US-10-310-002-3
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                                                                                                                           SEQ ID NO 106
LENGTH: 418
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APPLICANT: Emory University
TITLE OF INVENTION: Insulin-Responsive Sequence DNA Binding Protein-1 and Methods to
TITLE OF INVENTION: Regulate Insulin-Responsive Genes
TITLE OF INVENTION: Regulate Insulin-Responsive Genes
FILE REFERENCE: E056 10.0
CURRENT APPLICATION NUMBER: US/10/310,002
CURRENT FILING DATE: 2002-12-04
NUMBER OF SEQ ID NOS: 63
SOFTWARE: Patentin version 3.2
SEQ ID NO 47
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47.1%; Pred. No. 1.7e+03;
Live 5; Mismatches 3; Indels 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 57.1%; Pred. No. 1.9e+03;
Matches 8; Conservative 2; Mismatches 4; Indels C
4; Indels
  2; Mismatches
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CURRENT FILING DATE: 2002-05-29
PRIOR APPLICATION NUMBER: JP 2001-204089
PRIOR FILING DATE: 2001-06-30
PRIOR FILING DATE: 2001-06-30
PRIOR FILING DATE: 2001-08-02
NUMBER OF SEQ ID NOS: 15109
LENGTH: 793
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APPLICANT: SHIBA, TADAYOSHI
APPLICANT: SAKAKI, YOSHIVUKI
APPLICANT: HATTORI, MASAHIRA
TITLE OF INVENTION: NOVEL POLYNUCLEOTIDES
FILE REFERENCE: 249-262
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, Publication No. US20030119018A1
, GENERAL INFORMATION:
                                                                                                                                                                                                            ; Sequence 47, Application US/10310002; Publication No. US20030125296A1; GENERAL INFORMATION:
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ISHIKAWA, JUN
HORIKAWA, HIROSHI
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                                            2 LTVIAQQTŢSQGLL 15
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    8; Conservative
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hes 8; Conserv
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US-09-925-300-1641
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US-10-310-002-47
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TITLE OF INVENTION: Insulin-Responsive Sequence DNA Binding Protein-1 and Methods to
TITLE OF INVENTION: Regulate Insulin-Responsive Genes
FILE REFERENCE: E056 1010
CURRENT APPLICATION NUMBER: US/10/310,002
CURRENT FILING DATE: 2002-12-04
NUMBER OF SEQ ID NOS: 63
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                                                                                                                                                          Query Match
47.2%; Score 34; DB 10; Length 334;
Best Local Similarity 57.1%; Pred. No. 5e+02;
Matches 8; Conservative 1; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 50.0%; Pred. No. 8.1e+02;
Matches 7; Conservative 2; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: IKEDA, HAROON
APPLICANT: IKEDA, HAROON
APPLICANT: ISHIKAWA, JUN
APPLICANT: SHIBA, TADAYOSHI
APPLICANT: SHIBA, TADAYOSHI
APPLICANT: SHAKI, YOSHIYUKI
APPLICANT: HATTORI, WASAHIRA
TILE OF INVENTION: NOVEL POLYNUCLEOTIDES
FILE REFERENCE: 249-262
CURRENT APPLICATION NUMBER: US/10/156,761
CURRENT FILING DATE: 2002-05-29
PRIOR FILING DATE: 2001-05-30
PRIOR FILING DATE: 2001-05-30
PRIOR FILING DATE: 2001-05-30
PRIOR FILING DATE: 2001-05-30
PRIOR FILING DATE: 2001-05-30
PRIOR FILING DATE: 2001-05-30
PRIOR FILING DATE: 2001-05-30
PRIOR FILING DATE: 2001-05-30
                                                                                                                                                                                                                                                                                                                                                                                                                   , Sequence 12564, Application US/10156761
, Publication No. US20030119018A1
, GENERAL INFORMATION:
                                                                               ; ORGANISM: Corynebacterium glutamicum
US-09-738-626-5842
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US-10-310-002-11
Sequence 11, Application US/10310002
Publication No. US20030125296A1
GENERAL INFORMATION:
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US-10-156-761-12564
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SOFTWARE: Patentin ver. 3.0
SEQ ID NO 5842
LENGTH: 334
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: OMURA, SATOSHI APPLICANT: IKEDA, HARUO
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Best Local Similarity
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US-10-156-761-12564
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LENGTH: 508
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                                            LENGTH:
TYPE: PF
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45.8%; Score 33; DB 15; Length 102;
Best Local Similarity 60.0%; Pred. No. 1.9e+02;
Matches 6; Conservative 3; Mismatches 1; Indels 0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Gaps
                                         GENERAL INFORMATION:
APPLICANT: Craig Rosen,
APPLICANT: Craig Rosen,
TITLE OF INVENTOR Mucleic Acids, Proteins and Antibodies
FILE REPERENCE: PA101
CURRENT APPLICATION NUMBER: US/09/925,300
CURRENT FILING DATE: 2001-08.10
PRIOR FILING DATE: 2000-03-08
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 1890
SOFTWARE: Patentin Ver. 2.0
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 1641
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
45.8%; Score 33; DB 10; Length 81;
Best Local Similarity 43.8%; Pred. No. 1.5e+02;
Matches 7; Conservative 4; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: INEDA, HARUO
APPLICANT: IKEDA, HARUO
APPLICANT: IKEDA, HARUO
APPLICANT: ISHIKAWA, JUN
APPLICANT: BSHIKAWA, HIROSHI
APPLICANT: SHIBA, TADAYOSHI
APPLICANT: OSHIVINI
APPLICATION NUMBER: US/10/156,761
CURRENT APPLICATION NUMBER: UP 2001-204089
PRIOR APPLICATION NUMBER: UP 2001-204089
PRIOR APPLICATION NUMBER: UP 2001-205-29
PRIOR APPLICATION NUMBER: UP 2001-275697
PRIOR APPLICATION NUMBER: UP 2001-275697
NUMBER OF SEG ID NOS: 15109
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Tob time : 10.6248 secs
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Sequence 1641, Application US/09925300 Patent No. US20020151681A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT ORGANISM: Streptomyces avermitilis IS-10-156-761-12525
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 VLTVIAQQTTSQGLLS 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                           ; TYPE: PRT
; ORGANISM: Homo sapiens
JS-09-925-300-1641
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JS-10-156-761-12525
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